## 10306703.txt

## SEQUENCE LISTING

<110>	hort, Jay M. retz, Keith A. kray, Kevin A. karton, Nelson Robert karrett, James B. ' Donoghue, Eileen lathur, Eric J.
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Page 3

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                              200
Ser Cys Ser Leu Thr Gln Ala Leu Pro Ser Glu Leu Lys Val Ser Ala
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225 230 235 240
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## 10306703.txt

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catatog atg aaa gcg atc tta atc cca ttt tta tct ctt ctg att ccg
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1 5 10
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Leu Thr Pro Gln Ser Ala Phe Ala Gln Ser Glu Pro Glu Leu Lys Leu
gaa agt gtg gtg att gtc agt cgt cat ggt gtg cgt gct cca acc aag
Glu Ser Val Val Ile Val Ser Arg His Gly Val Arg Ala Pro Thr Lys
                                                                                           325
                                                                                           373
gcc acg caa ctg atg cag gat gtc acc cca gac gca tgg cca acc tgg
Ala Thr Gln Leu Met Gln Asp Val Thr Pro Asp Ala Trp Pro Thr Trp
ccg gta aaa ctg ggt tgg ctg aca ccg cgn ggt ggt gag cta atc gcc
Pro Val Lys Leu Gly Trp Leu Thr Pro Arg Gly Gly Glu Leu Ile Ala
65 70 75
                                                                                           421
                                                                                           469
tat ctc gga cat tac caa cgc cag cgt ctg gta gcc gac gga ttg ctg
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∠/10\ X	

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<213> Escherichia coli

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10306703.txt
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Asp Asn Val Ser Leu Thr Gly Ala Val Ser Leu Ala Ser Met Leu Thr
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Glu Ile Phe Leu Leu Gln Gln Ala Gln Gly Met Pro Glu Pro Gly Trp
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Gly Arg Ile Thr Asp Ser His Gln Trp Asn Thr Leu Leu Ser Leu His
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Asn Ala Gln Phe Tyr Leu Leu Gln Arg Thr Pro Glu Val Ala Arg Ser
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Arg Ala Thr Pro Leu Leu Asp Leu Ile Lys Thr Ala Leu Thr Pro His
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ccg caa tct gca ttc gct cag agt gag ccg gag ctg aag ctg gaa agt
Pro Gln Ser Ala Phe Ala Gln Ser Glu Pro Glu Leu Lys Leu Glu Ser
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Val Val Ile Val Ser Arg His Gly Val Arg Ala Pro Thr Lys Ala Thr
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Gln Leu Met Gln Asp Val Thr Pro Asp Ala Trp Pro Thr Trp Pro Val
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Lys 65	Leu	Gly	Glu	Leu	Thr 70	Pro	Arg	Gly		06703 G1u 75			Ala	Tyr	Leu 80		
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gac Asp 225	tgt Cys	gtc Val	tca Ser	tta Leu	acc Thr 230	ggt Gly	gcg Ala	gta Val	agc Ser	ctc Leu 235	gca Ala	tca Ser	atg Met	ctg Leu	acg Thr 240		720
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